- 1. An isolated nucleic acid molecule comprising a polynucleotide encoding a phospholipase $A_{2\gamma}$ polypeptide.
- 2. An isolated nucleic acid molecule according to claim 1, wherein said phospholipase $A_{2\gamma}$ polypeptide catalyzes cleavage of fatty acids from the *sn*-2-position of phospholipids.
- 3. An isolated nucleic acid molecule according to claim 2 wherein said polynuceotide encodes a sequence as set forth in SEQ ID NO:1 or SEQ ID NO:2.
 - 4. A vector comprising a nucleic acid molecule according to claim 1.
 - 5. A cell transformed or transfected with a vector according to claim 4.
- of An isolated nucleic acid molecule comprising a fragment of a polynucleotide encoding a phospholipase A2γ wherein said fragment specifically hybridizes with a sequence as set forth in SEQ ID NO:5 or SEQ ID NO:6.
- 7. An isolated nucleic acid comprising a polynucleotide having at least about 90% identity with SEQ ID NO:3, SEQ ID NO:4, SEQ ID NO:5, or SEQ ID NO:6.
- 8. An isolated nucleic acid according to claim 7 comprising SEQ ID NO:3, SEQ ID NO:4, SEQ ID NO:5, or SEQ ID NO:6.
- 9.—An antisense sequence which specifically hybridizes to SEQ ID NO:2, or SEQ ID NO:3.
 - 10. An isolated polypeptide comprising a phospholipase A2 γ .
- 11. An isolated polypeptide according to claim 10 which catalyzes cleavage of fatty acids from the *sn*-2-position of phospholipids.
 - 12. An isolated polypeptide according to claim 11 which has at least 90% identity with SEQ ID NO:1 or SEQ ID NO:2.
 - 13. An isolated polypeptide according to claim 12 comprising SEQ ID NO:1 or SEQ ID NO:2.
 - 14. An isolated polypeptide according to claim 12 which is a conservatively substituted variant of SEQ ID NO:1 or SEQ ID NO:2.
 - 15. An antibody capable of binding to a phospholipase $A2\gamma$ according to claim

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- 16. A method of treating inflammation in a patient, said method comprising decreasing calcium-independent phospholipase $A_2\gamma$ activity in the patient.
- 17. A method according to claim 16 wherein the patient suffers from Alzheimer's disease, myocardial ischemia, or myocardial infarction.
- 18. A method according to claim 17 comprising administering to the patient a phospholipase $A_2\gamma$ translational repressor molecule.
- 19. A method according to claim 17 comprising administering to the patient an antisense sequence which specifically hybridizes to SEQ ID NO:3 or SEQ ID NO:4.
- 20. A method of increasing fatty acid utilization in a patient in need thereof, said method comprising increasing iPLA₂ γ activity in the patient.
- 21. A method according to claim 20 wherein the patient suffers from diabetes or obesity.
- 22. A method according to claim 21 comprising administering to the patient a substance which blocks translational repression of $iPLA_2\gamma$ expression.
- 23. A method according to claim 21 comprising administering to the patient an $iPLA_2\gamma$ polypeptide as set forth in SEQ ID NO:1, SEQ ID NO:2 or a conservatively substituted variant thereof or administering a polynucleotide encoding said $iPLA_2\gamma$ polypeptide.
- 24. A method for measuring activity of a phospholipase $A_2\gamma$ polypeptide of cells in a biological sample, said method comprising introducing into the sample a phospholipid substrate for cleavage of fatty acids by the phospholipase $A_2\gamma$, wherein the phospholipase $A_2\gamma$ cleaves fatty acid from the *sn*-2-position of the phospholipid substrate and measuring cleavage of the phospholipid substrate, wherein measuring cleavage of the phospholipid substrate comprises quantifying the release of fatty acid from the substrate.
- 25. An assay method for identifying substances which modulate $iPLA_2\gamma$ expression in a cell, said method comprising contacting a candidate substance with cells comprising a promoter sequence operably linked to an $iPLA_2\gamma$ repressor binding site and a reporter gene and measuring expression of the reporter gene.
- 26. A method according to claim 25 wherein said repressor binding site comprises SEQ ID NO:7.

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- 27. A method according to claim 26 wherein said reporter gene encodes an enzyme capable of being detected by a colorimetric, fluorimetric or luminometric assay.
- 28. A method according to claim 27 wherein said reporter gene encodes a luciferase.
 - 29. A method according to claim 26 wherein the promoter sequence is a baculovirus promoter sequence.
 - 30. A method according to claim 26 wherein the cells are Sf9 cells.
 - 31. A genetically engineered cell capable of identifying substances which modulate iPLA_{2Y} expression in a cell, said cells comprising a promoter operably linked to an iPLA_{2Y} repressor binding site and a reporter gene.
 - 32. A genetically engineered cell according to claim 31 wherein said repressor binding site comprises SEQ ID NO:7.
 - 33. A genetically engineered cell according to claim 32 wherein said reporter gene encodes an enzyme capable of being detected by a colorimetric, fluorimetric or luminometric assay.
 - 34. A genetically engineered cell according to claim 33 wherein said reporter gene encodes a luciferase.
- 35. A genetically engineered cell according to claim 32 wherein the promoter is a baculovirus promoter.
- 36. A genetically engineered cell according to claim 32 wherein the cells are Sf9 cells.
- 37. A method for identifying a substance which modulates iPLA_{2Y} expression, the method comprising: (1) contacting a candidate substance with a repressor binding site and detecting binding to said site, or (2) contacting a candidate substance with cells capable of expressing iPLA_{2Y} or a fragment thereof and measuring the expression of iPLA_{2Y} or fragment thereof by the cells, wherein a level of expression greater or less than that in absence of the substance indicates activity in modulating iPLA_{2Y} expression.